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Human; secreted protein; EST; expressed sequence tag; diagnosis; forensic; gene therapy; chromosome mapping; signal peptide; upstream regulatory sequence; cytokine activity; cell proliferation; differentiation; haematopoiesis regulation; tissue growth regulation; reproductive hormone regulation; chemotactic; chemokinetic; haemostatic; thrombolytic; anti-inflammatory; tumour inhibition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX39440 to AAX39597 represent 5' expressed sequence tags (ESTs) for human secreted proteins, and encode the proteins given in AAX11374 to and an Viterminal fragment of a secreted protein. Siven the AAX11374 to and an Viterminal fragment of a secreted protein. The nucleic acid can also be used for producing secreted human gene products. They proteins obtained may have cytokine activity, cell proliferation/differentiation activity, haematopolesis regulating activity, tissue growth regulating activity, reproductive hormone regulating activity, chemostatic and
                                                                                                                                            61 SHPKTGTVDNNTSTDLKSLRPDELPHPEVDDLAQITTFWGQSPQTGCLPPDCSKCCHGDY 120
                                                                                                                                                                               SFRGYQGPPGPPGPPGIPGNHGNNGANGATGHEGAKGEKGDKGDLGPRGERGQHGPKGEK 107
                                                                                                                                                                                              New nucleic acids encoding human secreted - proteins obtained from cDNA libraries prepared from substantia nigra, cerebellum, surrenais and fetal brain tissue
                                                                                         -SPQTGGLPPDCSKCCHGDY
                                    Length 223;
                                                            Indels
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1e-68;
                             Score 792.5; D
Pred. No. 1e-68
0; Mismatches
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                                                                          1 MLWRQLIYWQLLALFFLPFCLCQDEYME------
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66.8%;
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 223 AA;
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thrombolytic activity, receptor/ ligand activity, anti-inflammatory activity, tumour inhibition activity or other activities. The products can be used in forensic, gene therapy and chromosome mapping procedures. The sequences can also be used for obtaining corresponding promoter sequences. The nucleic acids encoding the signal peptide can be used for directing extracellular secretion of a polypeptide or the insertion of a polypeptide into a membrane, or importing a polypeptide into a cell.
                                                                                                                                                                                                            Human; nootropic; immunosuppressant; cytostatic; gene therapy; cancer; peripheral nervous system; neuropathy; central nervous system; CNS; Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic; amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic; chemokinetic; thrombolytic; drug screening; arthritis; inflammation;
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                                                                                                                                                               1 MLWRQLIYWQLLALFFLPFCLCQDEYMESPQTGGLPPDCSKCCHGDYSFRGYGGPPGPPG
                                                                                                                                                                           Novel nucleic acids and polypeptides, useful for treating disorders such as central nervous system injuries -
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Zhang J;
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Yang Y,
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100.0%; Pred. No. 4.2e-62;
.ive 0; Mismatches 0;
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u C, Xue AJ,
Drmanac RT;
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Wang Z, Wehrman T, X
Zhou P, Goodrich R,
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2000US-0598042.
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                                                                                                                                             Conservative
                                                                                                                                Local Similarity
                                                                                             Sequence 128 AA;
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19-JUL-2000;
                                                                                                                                          Matches 124;
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Wang J, W
Zhao QA,
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GenCore version 5.1.3

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OM protein - protein search, using sw model

Run on: January 13, 2003, 10:42:48; Search time 15 Seconds
(without alignments)
482.536 Million cell updates/sec

Title: US-09-931-836-2
Perfect score: 1367
Sequence: 1 MLWRQLIXWQLLALFFLPFC......LHGDHQRFSTFAGFLLFETK 246
Scoring table: BLOSUM62
Gapop 10.0, Gapext.0.5
Searched: 262574 seqs, 29422922 residues
Total number of hits satisfying chosen parameters: 262574
Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Lissued_Patents_AA:*

1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Result		Query				
No.	Score	Match	Match Length	DB	ID	Description
н	582	42.6	105	. 4	[[S-09-188-930-147	
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APPLICANT: Waterson, James D.
APPLICANT: Strachan, Lorna
APPLICANT: Strachan, Matthew
APPLICANT: Sleeman, Matthew
APPLICANT: Ourst, Rene
APPLICANT: Murison, James Greg
APPLICANT: CANDON SCHOOL STREET STRE

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800011254507890012545	ALIGNM AESULT 1 S-09-188-330-147 Squence 147, Application US/09188930A Patent No. 6150502 GENERAL INFORMATION: APPLICANT: Waterson, James D. APPLICANT: Strachan, Matthew APPLICANT: Sleeman, Matthew APPLICANT: Sleeman, Matthew APPLICANT: Ornest, Rene APPLICANT: MINISON, James Greg TITLE OF INVENTION: Compositions Isolate TITLE OF INVENTION: Compositions Isolate TITLE OF INVENTION: AND MADBR: US/09/188, 9: CURRENT APPLICATION NUMBER: US/09/188, 9: CURRENT FILING DATE: 1998-11-09 NUMBER OF SEQ ID NOS: 346 SOFTWARE: FastSEQ for Windows Version 3 LENGTH: 105 TYPE: PRT ORGANISM: Rat	Ouery Match Best Local Similarity 93.3%; Pred. Matches 93.3%; Pred. Matches 93.3%; Pred. Matches 93.3%; Pred. Matches 94.3%; Pred. MIUMPOLIVEDIALFELPFCLCQDEYMESP 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Watson, James D.
APPLICANT: Strachan, Lorna
APPLICANT: Strachan, Lorna
APPLICANT: Strachan, Matthew
APPLICANT: Stream, Matthew
APPLICANT: Murison, James Greg
TITLE OF INVENTION: Compositions isolated From Skin Cells
TITLE OF INVENTION: and Methods For Their Use
FILE REFERENCE: 11000.1011c1
CURRENT APPLICATION NUMBER: US/09/188,930A
CURRENT FILING DATE: 1998-11-09
NUMBER OF SEQ ID NOS: 348
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23.1%; Score 316; DB 4; Length 294; 28.9%; Pred. No. 5.7e-24; ive 41; Mismatches 89; Indels
                                                                                                                                                 Length 105;
                                                                                                                                                                                                                                                                         61 PPGIPGNHGNNGANGATGHEGAKGEKGDKGDLGPRGERGQHGPKG 105
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Pred. No. 2.1e-51;
2; Mismatches 5
            NUMBER OF SEQ ID NOS: 348
SOFTWARE: FastSEQ for Windows Version 3.0
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CURRENT FILING DATE: 1998-11-09
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Best Local Similarity 93.3%;
Matches 98; Conservative
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Best Local Similarity 28.9%
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                                                                                            ; ORGANISM: Rat
US-09-188-930-280
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US-09-188-930-294
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LENGTH: 105
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Patent No. 6461821

GENERAL INFORMATION:

APPLICANT: Otsuka Pharmaceutical Co., Ltd.

TITLE OF INVENTION: Smooth muscle growth inhibitory composition, a

TITLE OF INVENTION: therefor

TITLE OF INVENTION: therefor

TITLE OF INVENTION: therefor

FILE REPERBUCE: P98-51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              34;
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                                                                                    NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
APPLICANT: Scherer, Philipp E.
APPLICANT: Lodish, Harvey F.
TITLE OF INVENTION: A NOVEL SERUM PROTEIN PRODUCED
TITLE OF INVENTION: EXCLUSIVELY IN ADIPOCYTES
                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/463,911
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 b; Score 294; DB 2; L
b; Pred. No. 7.4e-22;
40; Mismatches 102;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY AGENT INFORMATION:
NAME: Granhan, Patricia
REGISTATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: WH195-05
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 861-6240
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
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31.5%;
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amino acid
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Best Local Similarity 31.5%
Matches 81; Conservative
                                                                                                                                                                                                         Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: protein US-08-463-911-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 530
                                                                                                                                            STREET: Two
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                                                                                                                                                                                                                                                02173
                                                                                                                                                                                                                           COUNTRY:
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The invention relates to novel nucleic acid sequences derived from rat dermal papilla, human keratinocytes and neonatal foreskin fibroblasts, and mouse embryonic skin, Keratinocyte stem cells and tranalt amplifying calls. Polypeptides of the invention may be used to treat inflammation, cancer and neurological diseases. The proteins may be used to stimulate the growth and motility of keratinocytes, to inhibit the growth of cancer cells, to modulate skin inflammation, to modulate skin inflammation, to modulate epithelial cell growth and ovelopmental defects, skin wounds and hair follicle to treat growth and developmental defects, skin wounds and hair follicle to treat growth and developmental defects, skin wounds and hair follicle disorders. Sequences AAV75942-Y7613 represent polypeptides encoded by cDNA sequences AAV75942-Y7613 represent polypeptides encoded types. Sequences AAV75942-Y7513 represent polypeptides encoded by cDNA sequences AAV75942-Y75947, AAV7601-, AAV7601-Y7601, AAV
                                                  Claim 4; Page 112-113; 235pp; English.
         including wounds and cancer
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                                    The invention relates to human nucleic acids (AAL57798-AAI61369) and the encoded polypeptides (AAM38642-AAM42213) with nootropic, immunosuppressant and cytostatic activity. The polynucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous system, such as peripheral nervous injuries, peripheral nervous alteral nervous system diseases, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activities such as Immune system suppression and thrombolytic activity, chemctactic/chemokinetic activity, hemostatic and thrombolytic activity, arthritis and inflammation, leukaemias and
                                                                                                                                                                                                                                                                                                                                                                     The sequence data for this patent did not form part of the printed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Skin; dermal papilla; keratinocyte; neonatal foreskin fibroblast; embryonic skin cell; keratinocyte stem cell; transit amplifying cell; secreted; transmembrane; infibammation; cancer; neurological disease; anglogenesis; tumour vascularisation; growth disorder; developmental disorder; skin wound; hair follicle disorder; anti-inflammatory; cytostatic; neuroprotective; vulnery.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              121 MASLATHFSNQNSCIIFSSVETNIGNFFDVMTGRFGAPVSGVYFFTFSMMKHEDVEEVYV 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Pred. No. 1.8e-56;
0; Mismatches 0;
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Example 5; SEQ ID NO 3219; 10078pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                         126 AA;
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|121 LLFETK 126
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                                                                                                                                                                                                                                                                                                                                                                                           specification
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09-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Murison JG;
                                                                                                               1 MIWRQLIYWQLLALFFLPFCLCQDEYMESPQTGGLPPDCSKCCHGDYSFRGYGGPPGPPG 60
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                                                                         Indels · 0;
                                              Score 582; DB 21; Length 105; Pred. No. 1e-48; 2; Mismatches 5; Indels · (
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                                                                                                                                                                  61 PPGIPGNHGNNGANGATGHEGAKGEKGDKGDLGPRGERGQHGPKG 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Onrust R,
                                                                                                                                                                                                                                                                                                                        Rat skin cell protein, SEQ ID NO:280.
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putative transmembrane domains.
                                               42.6%;
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                                                                         Conservative
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                                                         Local Similarity
les 98; Conservat
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                       105 AA;
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09-NOV-1998;
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                       Sequence
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                                              Query Match
                                                                         Matches
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Search completed: January 13, 2003, 10:42:49
Job time : 38 secs
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                                                The invention relates to novel nucleic acid sequences derived from rat dermal papilla, human keratinocytes and neonatal foreskin fibroblasts, and mouse embryonic skin, keratinocyte stem cells and transit amplifying cells. Polypeptides of the invention may be used to treat inflammation, the growth and motility of keratinocytes, to inhibit the growth of cancer cells, to modulate angiogenesis and tumour vascularisation, to modulate skin inflammation, to modulate skin inflammation, to modulate epithelial cell growth and to inhibit binding of HIV-1 to leukocytes. The invention may also be used to treat growth and developmental defects, skin wounds and hair follicle disorders. Sequences AAY75942-Y75947, AAY76012. Tepresent polypeptides encoded by CDNA sequences AAY75942-Y75947, AAY76011, AAY76014-Y76104 and AAY76119 are proteins with an N-terminal signal sequence, indicating that they are secreted. Sequences AAY75986-Y75989, AAY76019 and AAY76119.
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                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                        1 MLWRQLIYWQLLALFFLPFCLCQDEYMESPQTGGLPPDCSKCCHGDYSFRGYQGPPGPFG 60
                                                                                                                                                                                                                                                                                                                                                                                                           New isolated polynucleotide used in the identification of genetic disorders and encoding polypeptides used for treating inflammatory disease, cancer and neurological diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rat; skin cell; cytostatic; antiinflammatory; anti-HIV; nootropic; neuroprotective; vulnerary; immunomodulatory; vaccine; keratinocyte growth stimulation; cancer; anglogenesis inhibition; inflammation; neurological disease.
                                                                                                                                                                                                                                                                                                                                    Score 582; DB 21; Length 105;
Pred. No. 1e-48;
2; Mismatches 5; Indels
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                          Page 168; 235pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Skin cell protein, SEQ ID NO: 147.
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                                                                                                                                                                                                                                                                             putative transmembrane domains
                                                                                                                                                                                                                                                                                                                                    42.6%;
 and cancer
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N-PSDB; AAC99566, AAC99776.
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Matches 98; Conservative
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                                                                                                                                                                                                                                                                                                          105 AA;
 including wounds
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                                                 The present sequence is a polypeptide which is expressed in mammalian skin cells. The polypeptide is useful for stimulating keratinocyte growth and motility, inhibiting the growth of cancer cells, modulating angiogenesis, inhibiting angiogenesis and vascularisation of tumours, modulating skin inflammation, stimulating the growth of epithelial cells, inhibiting the binding of human immunodeficiency virus (HIV)-1 to leukocytes, and treating inflammatory disease, cancer and neurological diseases. The polynucleotide can be used as a marker, in the identification of genetic disorders, and for the design of oligonucleotides for examining expression patterns.
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 582; DB 22
Pred. No. 1e-48;
2; Mismatches
Claim 4; Page 147; 352pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 93.3%;
Matches 98; Conservative
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